

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: 10/559,952  
Source: 1FWP  
Date Processed by STIC: 12/20/05

# ***ENTERED***



IFWP

## RAW SEQUENCE LISTING

DATE: 12/20/2005

PATENT APPLICATION: US/10/559,952

TIME: 10:25:16

Input Set : A:\1003 SEQLIST.txt

Output Set: N:\CRF4\12202005\J559952.raw

4 <110> APPLICANT: Roche, Andrew  
 5 Hansen, Martin Chr.  
 6 Villsen, Inge D.  
 7 Schrotz-King, Petra  
 8 Henningsen, Jeannette  
 9 Lund Jorgensen, Trine Louise  
 11 <120> TITLE OF INVENTION: Extracellular Aspergillus Polypeptides  
 14 <130> FILE REFERENCE: 13403.1003  
 C--> 16 <140> CURRENT APPLICATION NUMBER: US/10/559,952  
 C--> 16 <141> CURRENT FILING DATE: 2005-12-09  
 16 <160> NUMBER OF SEQ ID NOS: 49  
 18 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
 20 <210> SEQ ID NO: 1  
 21 <211> LENGTH: 260  
 22 <212> TYPE: PRT  
 23 <213> ORGANISM: Aspergillus Fumigatus  
 25 <400> SEQUENCE: 1  
 26 Met Leu Ala Ser Phe Gln Phe Cys Ile Leu Pro Arg Thr Tyr Arg Thr  
 27 1 5 10 15  
 28 Leu Leu Cys Ser Ala Gly Ala Gly Pro Leu Leu Ile Ile Gln Phe Val  
 29 20 25 30  
 30 Thr Val Ala Ser Ala Leu Ala Leu Ala Pro Thr Ala Val Val Ala Arg  
 31 35 40 45  
 32 Gln Gly Ala Ala Ala Phe Val Thr Val Asn Ser Ile Asp Val Cys Pro  
 33 50 55 60  
 34 Lys Lys Val Ala Gln Glu Ile Ile Asn Pro Gly Pro Lys Val Val Thr  
 35 65 70 75 80  
 36 Thr Pro Tyr Thr Cys Asp Gln Val Lys Leu Gly His Gly Leu Asp Val  
 37 85 90 95  
 38 Ser Tyr Tyr Asn Phe Asp Ile Glu Pro Leu Thr Lys Asp Thr Phe Pro  
 39 100 105 110  
 40 Tyr Cys Lys Ala Leu Lys Val Phe Asp Asn Glu Gly Cys Leu Gly Phe  
 41 115 120 125  
 42 Pro Thr Leu Trp Ile Pro Leu Glu Ser Pro Leu Glu Asp Lys Cys Ile  
 43 130 135 140  
 44 Pro Glu His Tyr Phe Ser Asp Glu Val Lys Ser Ile Ser Phe Gln Leu  
 45 145 150 155 160  
 46 Asp Cys Arg Glu Asp Ala Pro Val Lys Lys Glu Pro Tyr Gly Pro Lys  
 47 165 170 175  
 48 Glu Gly Ala Glu Gln Ser Ala Pro Gln Ala Glu His Ser Thr Lys Gln  
 49 180 185 190  
 50 Asp Ala Gln Gln Gly Ser His Gln Gly Gln Glu Val Gln Asn Ser Pro  
 51 195 200 205

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52 Lys Gln Glu Ala Arg Gln Gly Ser Arg Pro Ala Glu Ala Ala Pro Lys
53      210                      215                      220
54 Gln Glu Gln Glu Ala Glu Gln Ala Ser Glu Ala Ala Pro Glu Lys Lys
55 225                      230                      235                      240
56 Ala Ser Asn Pro Ala Asp Ser Leu Gly Leu Gly Glu Leu Thr Lys Val
58                      245                      250                      255
59 Leu Gly Phe Arg
60      260
62 <210> SEQ ID NO: 2
63 <211> LENGTH: 107
64 <212> TYPE: PRT
65 <213> ORGANISM: Aspergillus Fumigatus
67 <400> SEQUENCE: 2
68 Val Arg Phe Pro Val Pro Asp Asp Ile Thr Val Lys Gln Ala Thr Glu
69 1      5                      10                      15
70 Lys Cys Gly Asp Gln Ala Gln Leu Ser Cys Cys Asn Lys Ala Thr Tyr
71      20                      25                      30
72 Ala Gly Asp Val Thr Asp Ile Asp Glu Gly Ile Leu Ala Gly Thr Leu
73      35                      40                      45
74 Lys Asn Leu Ile Gly Gly Gly Ser Gly Thr Glu Gly Leu Gly Leu Phe
75      50                      55                      60
76 Asn Gln Cys Ser Lys Leu Asp Leu Gln Ser Pro Ile Ile Gly Ile Pro
77 65                      70                      75                      80
78 Ile Gln Asp Leu Val Asn Gln Lys Cys Lys Gln Asn Ile Ala Cys Cys
79      85                      90                      95
80 Gln Asn Ser Pro Ser Asp Ala Val Arg Phe Pro
81      100                      105
83 <210> SEQ ID NO: 3
84 <211> LENGTH: 318
85 <212> TYPE: PRT
86 <213> ORGANISM: Aspergillus Fumigatus
88 <400> SEQUENCE: 3
89 Met Ala Thr Pro Lys Val Gly Ile Asn Gly Phe Gly Arg Ile Gly Arg
90 1      5                      10                      15
91 Ile Val Gly Leu Asn Ser Leu Ser His Gly Val Asp Val Val Ala Val
92      20                      25                      30
93 Asn Asp Pro Phe Ile Glu Val His Tyr Ala Ala Tyr Met Leu Lys Tyr
94      35                      40                      45
95 Asp Thr Thr His Gly Gln Phe Lys Gly Thr Ile Glu Thr Tyr Asp Gln
96      50                      55                      60
97 Gly Leu Ile Val Asn Gly Lys Lys Ile Arg Phe Tyr Ala Glu Lys Asp
98 65                      70                      75                      80
99 Pro Ser Gln Ile Pro Trp Ser Glu Thr Gly Ala Ala Tyr Ile Val Glu
100      85                      90                      95
101 Ser Thr Gly Val Phe Thr Thr Lys Glu Lys Ala Ser Ala His Leu Lys
102      100                      105                      110
103 Gly Gly Ala Lys Lys Val Ile Ile Ser Ala Pro Ser Ala Asp Ala Pro
104      115                      120                      125
105 Met Phe Val Met Gly Val Asn Asn Thr Thr Tyr Thr Ser Asp Ile Gln

```

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```

106      130      135      140
107 Val Leu Ser Asn Ala Ser Cys Thr Thr Asn Cys Leu Ala Pro Leu Ala
108 145      150      155      160
109 Lys Val Ile Asn Asp Lys Phe Gly Ile Val Glu Gly Leu Met Thr Thr
110      165      170      175
111 Val His Ser Tyr Thr Ala Thr Gln Lys Val Val Asp Ala Pro Ser Asn
112      180      185      190
113 Lys Asp Trp Arg Gly Gly Arg Thr Ala Ala Gln Asn Ile Ile Pro Ser
114      195      200      205
115 Ser Thr Gly Ala Ala Lys Ala Val Gly Lys Val Ile Pro Ser Leu Asn
116      210      215      220
117 Gly Lys Leu Thr Gly Met Ala Met Arg Val Pro Thr Ser Asn Val Ser
118 225      230      235      240
119 Val Val Asp Leu Thr Cys Arg Leu Glu Lys Gly Ala Ser Tyr Asp Glu
120      245      250      255
121 Ile Lys Gln Ala Ile Lys Ala Ala Ser Glu Glu Gly Glu Leu Lys Asn
122      260      265      270
123 Ile Leu Gly Tyr Thr Glu Asp Asp Val Val Ser Ser Asp Leu Asn Gly
124      275      280      285
125 Asp Glu Arg Ser Ser Ile Phe Asp Ala Lys Ala Gly Ile Ser Leu Asn
126      290      295      300
127 Pro Asn Phe Val Lys Leu Val Ala Trp Tyr Asp Asn Glu Trp
128 305      310      315
130 <210> SEQ ID NO: 4
131 <211> LENGTH: 438
132 <212> TYPE: PRT
133 <213> ORGANISM: Aspergillus Fumigatus
135 <400> SEQUENCE: 4
136 Met Pro Ile Ser Lys Ile His Ala Arg Ser Val Tyr Asp Ser Arg Gly
137 1      5      10      15
138 Asn Pro Thr Val Glu Val Asp Val Ala Thr Glu Thr Gly Leu His Arg
139      20      25      30
140 Ala Ile Val Pro Ser Gly Ala Ser Thr Gly Gln His Glu Ala His Glu
141      35      40      45
142 Leu Arg Asp Gly Asp Lys Thr Gln Trp Gly Gly Lys Gly Val Leu Lys
143      50      55      60
144 Ala Val Lys Asn Val Asn Glu Thr Ile Gly Pro Ala Leu Ile Lys Glu
145 65      70      75      80
146 Asn Ile Asp Val Lys Asp Gln Ser Lys Val Asp Glu Phe Leu Asn Lys
147      85      90      95
148 Leu Asp Gly Thr Ala Asn Lys Ser Asn Leu Gly Ala Asn Ala Ile Leu
149      100      105      110
150 Gly Val Ser Leu Ala Val Ala Lys Ala Gly Ala Ala Glu Lys Gly Val
151      115      120      125
152 Pro Leu Tyr Ala His Ile Ser Asp Leu Ala Gly Thr Lys Lys Pro Tyr
153      130      135      140
154 Val Leu Pro Val Pro Phe Gln Asn Val Leu Asn Gly Gly Ser His Ala
155 145      150      155      160
156 Gly Gly Arg Leu Ala Phe Gln Glu Phe Met Ile Val Pro Asp Ser Ala

```

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```

157          165          170          175
158 Pro Ser Phe Ser Glu Ala Leu Arg Gln Gly Ala Glu Val Tyr Gln Lys
159          180          185          190
160 Leu Lys Ala Leu Ala Lys Lys Lys Tyr Gly Gln Ser Ala Gly Asn Val
161          195          200          205
162 Gly Asp Glu Gly Gly Val Ala Pro Asp Ile Gln Thr Ala Glu Glu Ala
163          210          215          220
164 Leu Asp Leu Ile Thr Glu Ala Ile Glu Gln Ala Gly Tyr Thr Gly Lys
165 225          230          235          240
166 Ile Lys Ile Ala Met Asp Val Ala Ser Ser Glu Phe Tyr Lys Ala Asp
167          245          250          255
168 Val Lys Lys Tyr Asp Leu Asp Phe Lys Asn Pro Glu Ser Asp Pro Ser
169          260          265          270
170 Lys Trp Leu Thr Tyr Glu Gln Leu Ala Asp Leu Tyr Lys Ser Leu Ala
171          275          280          285
172 Ala Lys Tyr Pro Ile Val Ser Ile Glu Asp Pro Phe Ala Glu Asp Asp
173          290          295          300
174 Trp Glu Ala Trp Ser Tyr Phe Tyr Lys Thr Ser Asp Phe Gln Ile Val
175 305          310          315          320
176 Gly Asp Asp Leu Thr Val Thr Asn Pro Gly Arg Ile Lys Lys Ala Ile
177          325          330          335
178 Glu Leu Lys Ser Cys Asn Ala Leu Leu Leu Lys Val Asn Gln Ile Gly
179          340          345          350
180 Thr Leu Thr Glu Ser Ile Gln Ala Ala Lys Asp Ser Tyr Ala Asp Asn
181          355          360          365
182 Trp Gly Val Met Val Ser His Arg Ser Gly Glu Thr Glu Asp Val Thr
183          370          375          380
184 Ile Ala Asp Ile Ala Val Gly Leu Arg Ser Gly Gln Ile Lys Thr Gly
185 385          390          395          400
186 Ala Pro Cys Arg Ser Glu Arg Leu Ala Lys Leu Asn Gln Ile Leu Arg
187          405          410          415
188 Ile Glu Glu Glu Leu Gly Glu Asn Thr Val Tyr Ala Gly Ser Lys Phe
189          420          425          430
190 Arg Thr Ala Val Asn Leu
191          435
193 <210> SEQ ID NO: 5
194 <211> LENGTH: 728
195 <212> TYPE: PRT
196 <213> ORGANISM: Aspergillus Fumigatus
198 <400> SEQUENCE: 5
199 Met Arg Leu Thr Phe Ile Pro Ser Leu Ile Gly Val Ala Asn Ala Val
200 1          5          10          15
201 Cys Pro Tyr Met Thr Gly Glu Leu Asn Arg Arg Asp Glu Ile Ser Asp
202          20          25          30
203 Gly Asp Ala Ala Ala Thr Glu Glu Phe Leu Ser Gln Tyr Tyr Leu
204          35          40          45
205 Asn Asp Asn Asp Ala Phe Met Thr Ser Asp Val Gly Gly Pro Ile Glu
206          50          55          60
207 Asp Gln Asn Ser Leu Ser Ala Gly Glu Arg Gly Pro Thr Leu Leu Glu

```

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Input Set : A:\1003 SEQLIST.txt

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208	65					70				75				80		
209	Asp	Phe	Ile	Phe	Arg	Gln	Lys	Ile	Gln	Arg	Phe	Asp	His	Glu	Arg	Val
210					85					90				95		
211	Pro	Glu	Arg	Ala	Val	His	Ala	Arg	Gly	Ala	Gly	Ala	His	Gly	Val	Phe
212				100					105					110		
213	Thr	Ser	Tyr	Gly	Asp	Phe	Ser	Asn	Ile	Thr	Ala	Ala	Ser	Phe	Leu	Ala
214			115					120					125			
215	Lys	Glu	Gly	Lys	Gln	Thr	Pro	Val	Phe	Val	Arg	Phe	Ser	Thr	Val	Ala
216		130					135					140				
217	Gly	Ser	Arg	Gly	Ser	Ser	Asp	Leu	Ala	Arg	Asp	Val	His	Gly	Phe	Ala
218	145				150					155					160	
219	Thr	Arg	Phe	Tyr	Thr	Asp	Glu	Gly	Asn	Phe	Asp	Ile	Val	Gly	Asn	Asn
220				165					170					175		
221	Ile	Pro	Val	Phe	Phe	Ile	Gln	Asp	Ala	Ile	Leu	Phe	Pro	Asp	Leu	Ile
222			180					185					190			
223	His	Ala	Val	Lys	Pro	Arg	Gly	Asp	Asn	Glu	Ile	Pro	Gln	Ala	Ala	Thr
224		195					200					205				
225	Ala	His	Asp	Ser	Ala	Trp	Asp	Phe	Phe	Ser	Gln	Gln	Pro	Ser	Thr	Met
226		210					215				220					
227	His	Thr	Leu	Leu	Trp	Ala	Met	Ser	Gly	His	Gly	Ile	Pro	Arg	Ser	Phe
228	225				230					235					240	
229	Arg	His	Val	Asp	Gly	Phe	Gly	Val	His	Thr	Phe	Arg	Phe	Val	Thr	Asp
230				245					250					255		
231	Asp	Gly	Ala	Ser	Lys	Leu	Val	Lys	Phe	His	Trp	Lys	Ser	Leu	Gln	Gly
232			260					265					270			
233	Lys	Ala	Ser	Met	Val	Trp	Glu	Glu	Ala	Gln	Gln	Thr	Ser	Gly	Lys	Asn
234		275					280				285					
235	Pro	Asp	Phe	Met	Arg	Gln	Asp	Leu	His	Asp	Ala	Ile	Glu	Ala	Gly	Arg
236		290				295				300						
237	Tyr	Pro	Glu	Trp	Glu	Leu	Gly	Val	Gln	Ile	Met	Asp	Glu	Glu	Asp	Gln
238	305				310					315				320		
239	Leu	Arg	Phe	Gly	Phe	Asp	Leu	Leu	Asp	Pro	Thr	Lys	Ile	Val	Pro	Glu
240				325					330					335		
241	Glu	Phe	Val	Pro	Ile	Thr	Lys	Leu	Gly	Lys	Met	Gln	Leu	Asn	Arg	Asn
242			340					345					350			
243	Pro	Arg	Asn	Tyr	Phe	Ala	Glu	Thr	Glu	Gln	Val	Met	Phe	Gln	Pro	Gly
244		355					360					365				
245	His	Ile	Val	Arg	Gly	Val	Asp	Phe	Thr	Glu	Asp	Pro	Leu	Leu	Gln	Gly
246		370				375					380					
247	Arg	Leu	Phe	Ser	Tyr	Leu	Asp	Thr	Gln	Leu	Asn	Arg	His	Gly	Gly	Pro
248	385				390				395					400		
249	Asn	Phe	Glu	Gln	Leu	Pro	Ile	Asn	Gln	Pro	Arg	Val	Pro	Val	His	Asn
250				405					410					415		
251	Asn	Asn	Arg	Asp	Gly	Ala	Gly	Gln	Met	Phe	Ile	Pro	Leu	Asn	Pro	His
252			420					425					430			
253	Ala	Tyr	Ser	Pro	Lys	Thr	Ser	Val	Asn	Gly	Ser	Pro	Lys	Gln	Ala	Asn
254		435					440					445				
255	Gln	Thr	Val	Gly	Asp	Gly	Phe	Phe	Thr	Ala	Pro	Gly	Arg	Thr	Thr	Ser
256		450				455					460					

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/559,952

DATE: 12/20/2005  
TIME: 10:25:17

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:36; Xaa Pos. 176,179

Seq#:37; Xaa Pos. 2,5

Seq#:38; N Pos. 579,580,581,588,589,590

Seq#:39; N Pos. 526,527,528,535,536,537

## VERIFICATION SUMMARY

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PATENT APPLICATION: US/10/559,952

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Input Set : A:\1003 SEQLIST.txt

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L:16 M:270 C: Current Application Number differs, Replaced Current Application No  
L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:722 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:160  
L:724 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:176  
L:760 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0  
L:783 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:540  
L:815 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:480